

1/42

10/539565

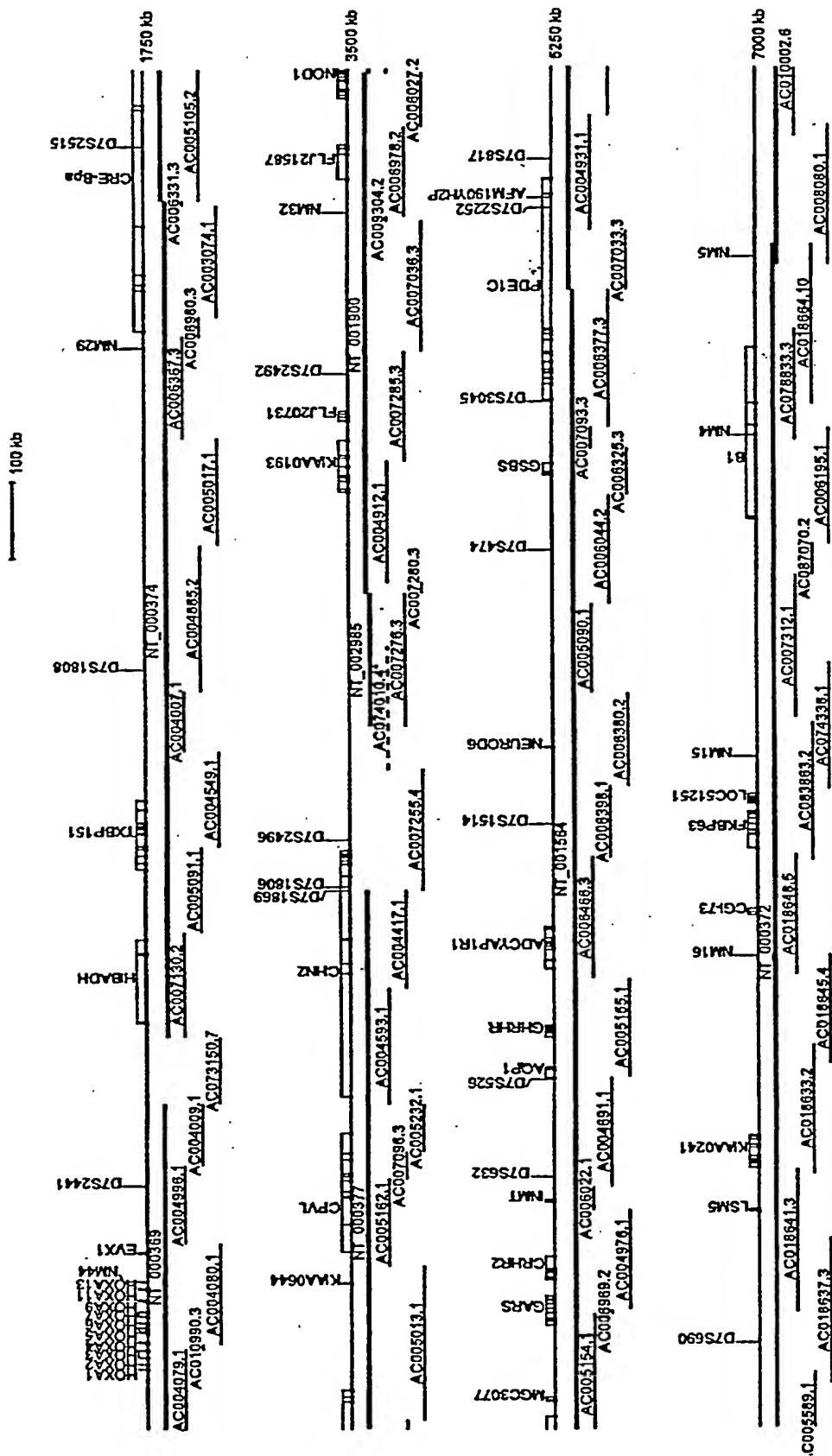


Fig. 1

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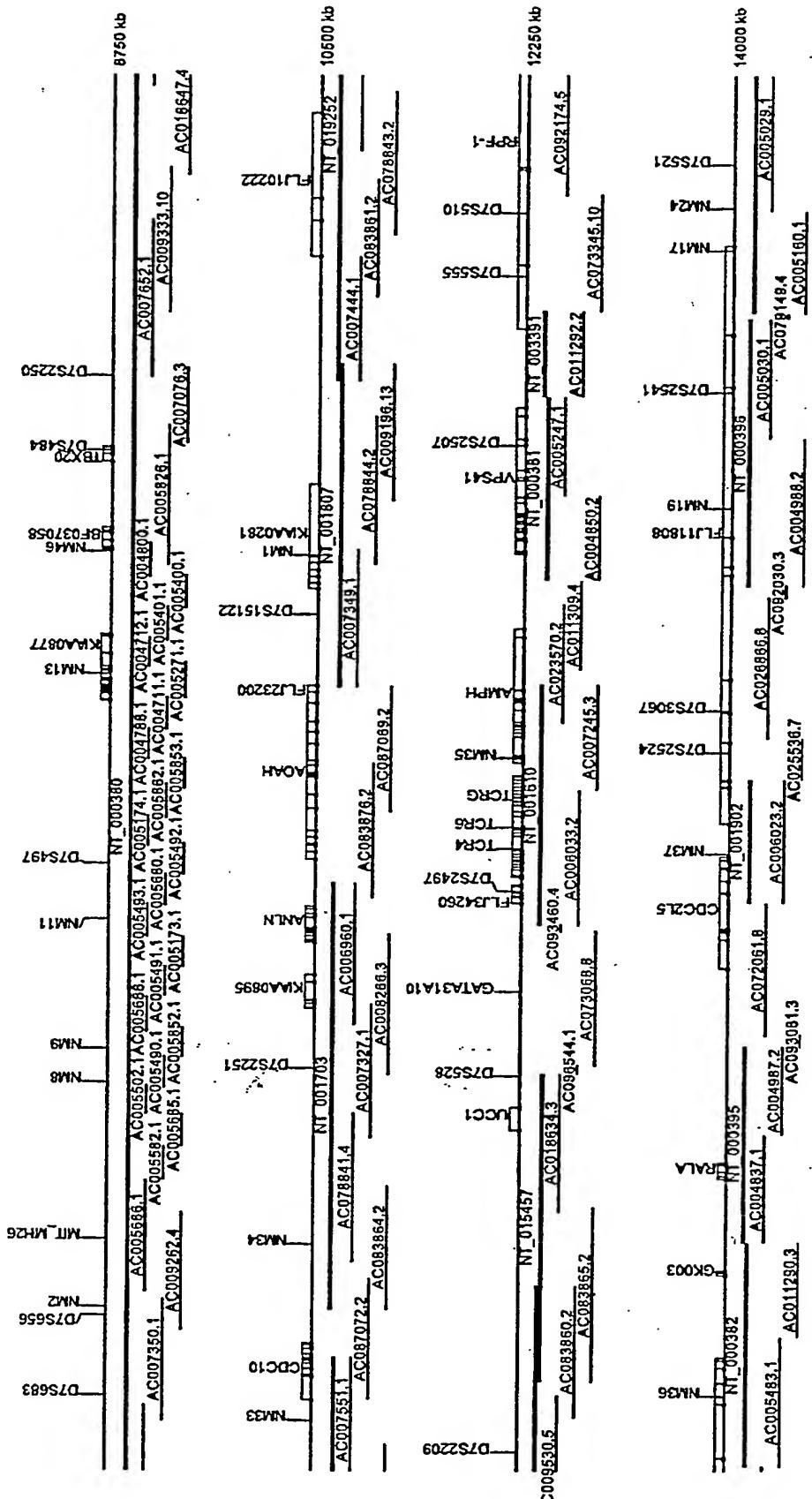


Fig. 1 (continued)

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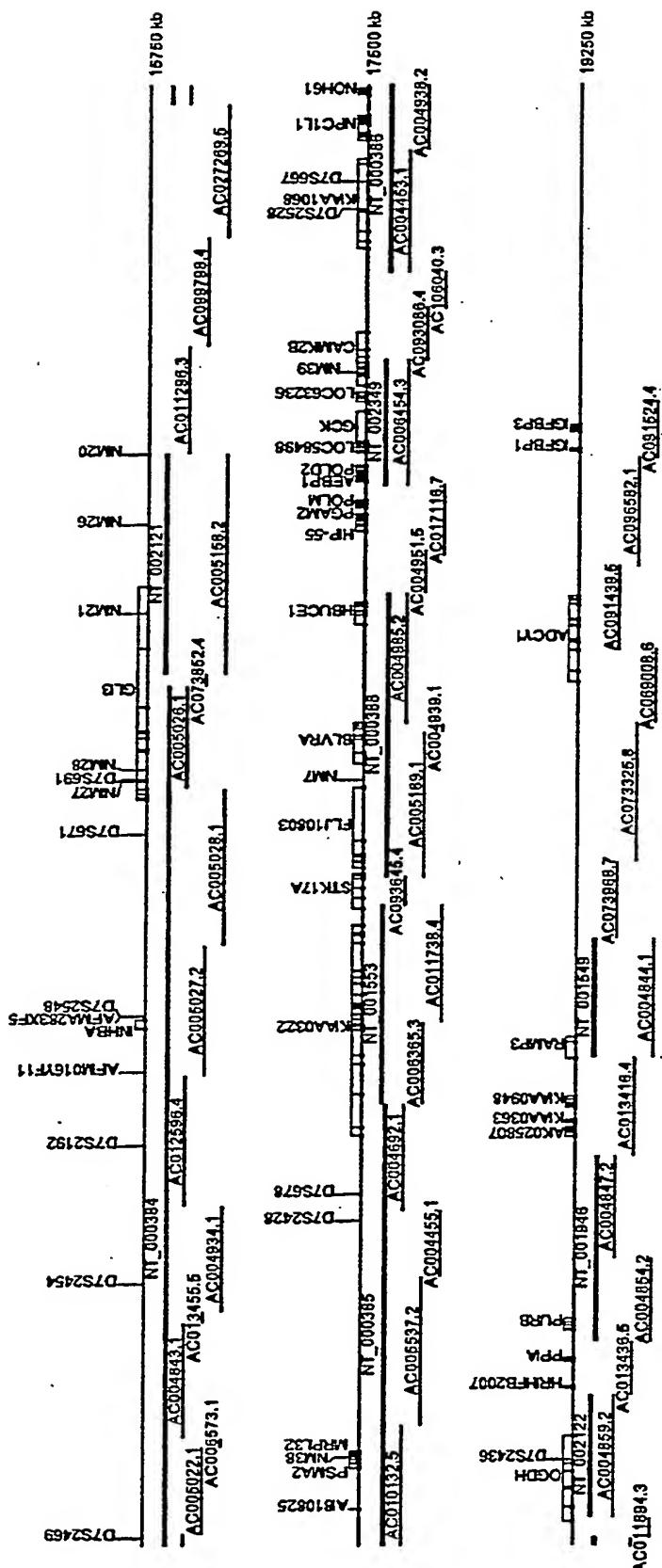


Fig. 1 (continued)

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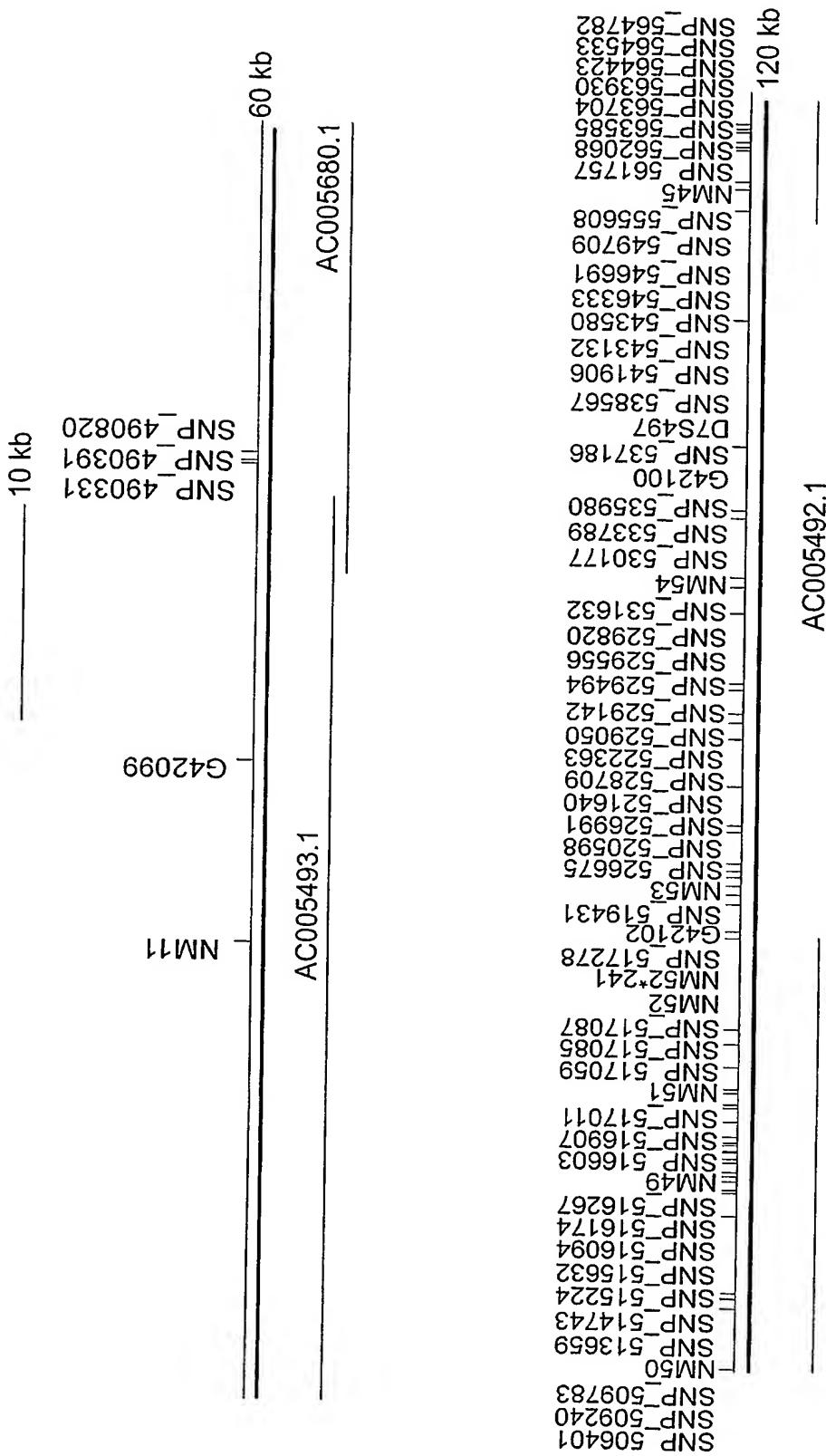


Fig. 2

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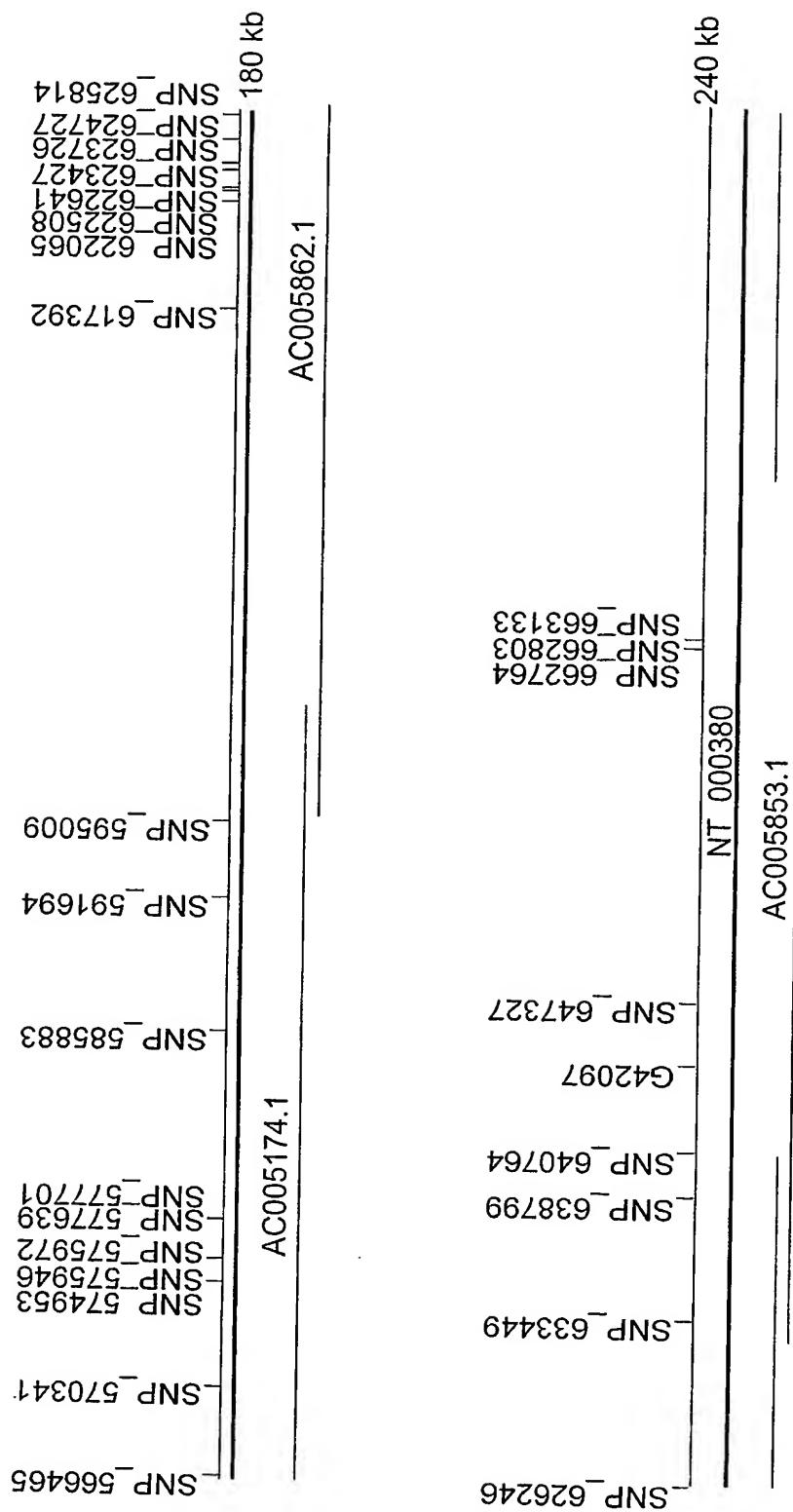


Fig. 2 (Continued)

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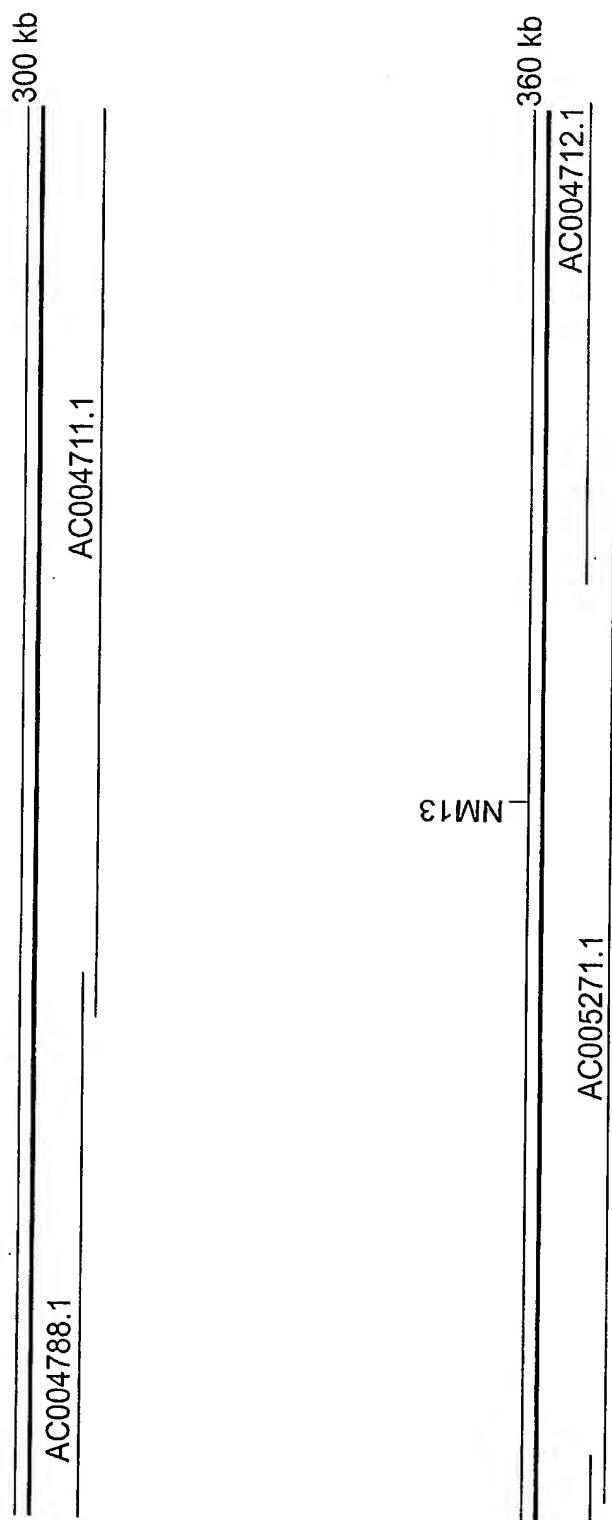


Fig. 2 (Continued)

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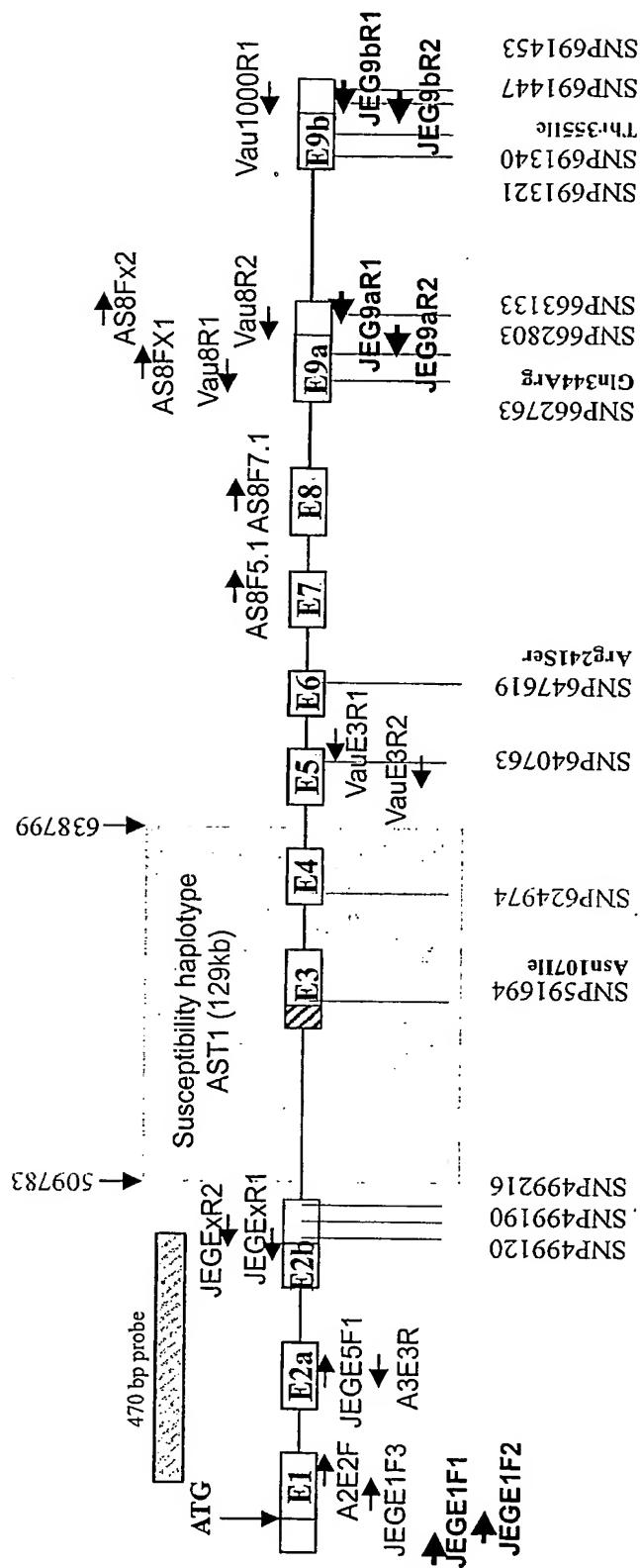


Fig. 3

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Figure 4A

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Figure 4B1

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Figure 4C

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Figure 4D

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Figure 4E

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Figure 4F

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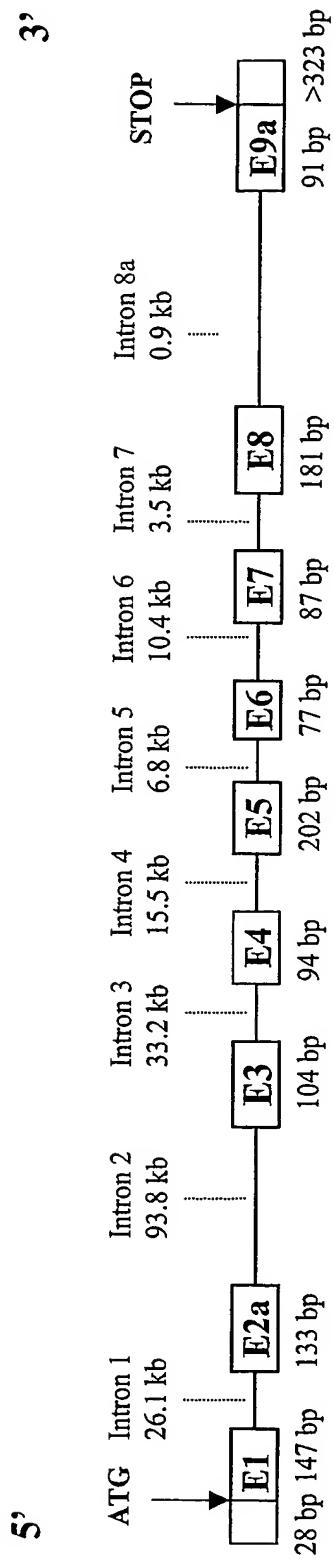


Figure 5A

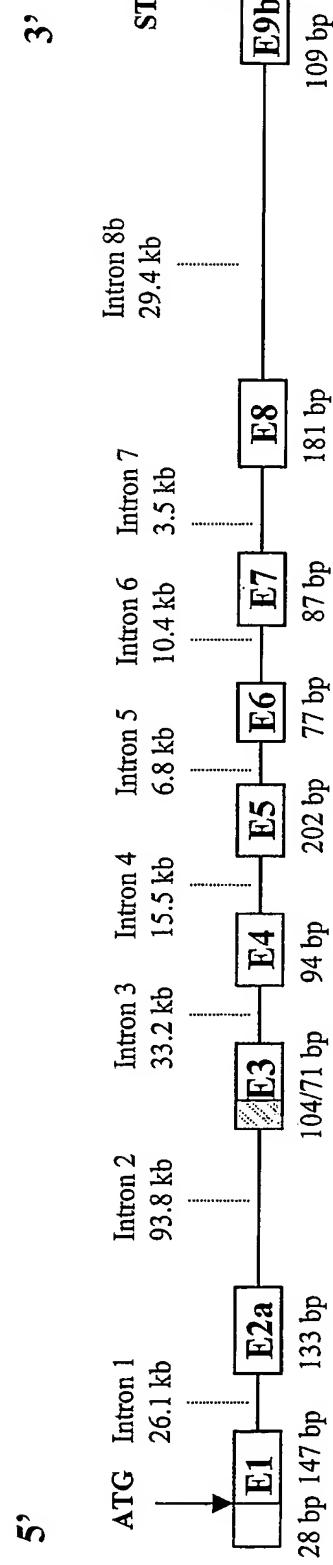


Figure 5B

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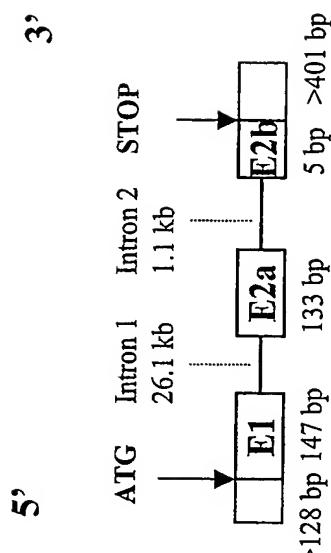


Figure 5C

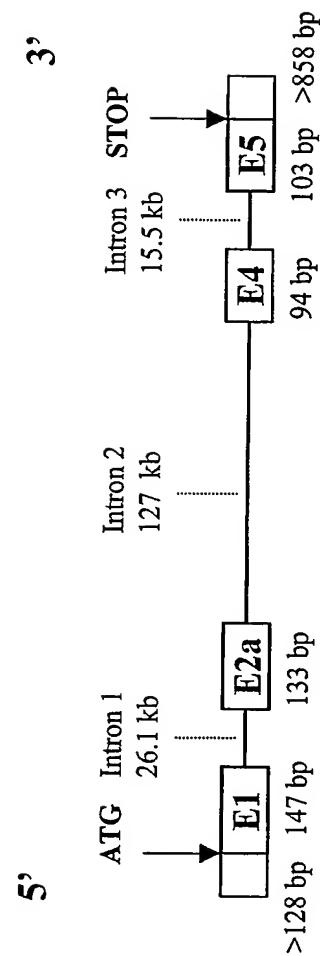


Figure 5D

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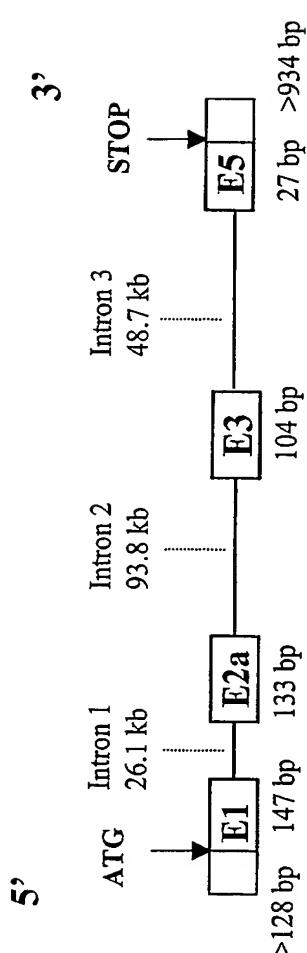


Figure 5E

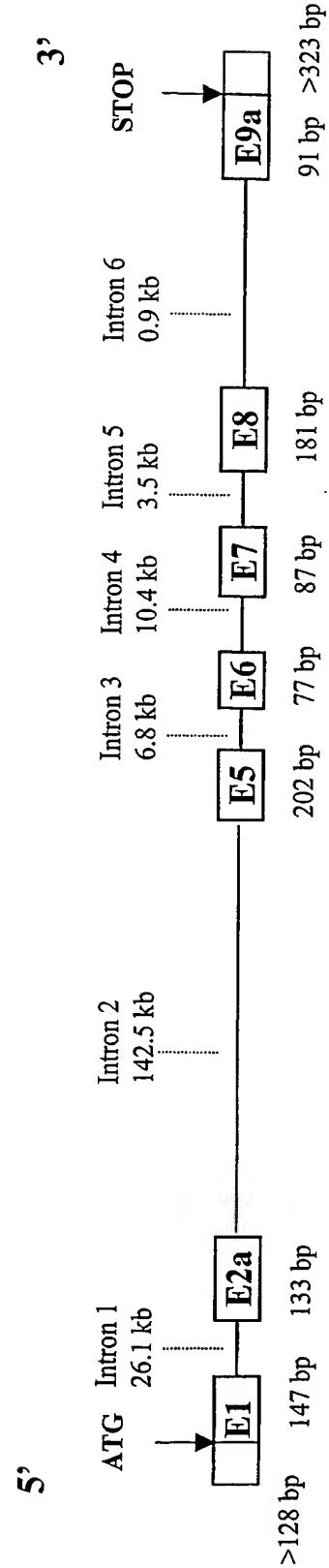


Figure 5F

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MPANFTEGSFDSSGTGQTLDSSPVACTETVTFTEVVEGKEWGSFYYSFKT

TM1 CYTOLOOP1 TM2
EQLITLWVLFVFTIVGNNSVVLFSTWRRKKSRMTFFVTQLAITDSFTGLV

EXOLOOP1 TM3
NILTDINWRFTGDFTAPDLVCRVVRYLQVVLLYASTYVLVSLSIDRYHAI

CYTOLOOP2 TM4 EXOLOOP2
VYPMKFLQGEKQARVLIVIAWSLSFLFSIPTLIIFGKRTLSNGEVQCWAL

TM5 CYTOLOOP3
WPDDSYWTPYMTIVAFLVYFIPLTIISIMYGIVIRTIWIKSKTYETVISN

TM6
CSDGKLCSSYNRGLISKAKIKAIKYSSIIILAFICCWSPYFLFDILDNFN

EXOLOOP3 TM7
LLPDTQERFYASVIIQNLPALNSAINPLIYCVFSSSISFPCREQRSQDSR

MTFRERTERHEMQILSKPEFI*

Figure 6A

101539565

B_{long}

MPANFTEGSFDSSGTGQTLDSVPACTETVTFTEVVEGKEWGSFYYSFKT

TM1 **CYTOLOOP1** **TM2**
EQLITLWVLVFVFTIVGNSVVLFSTWRRKKKSRTFFVTQLAITDSFTGLV

EXOLOOP1 **TM3**
NILTDINWRFTGDFTAPDLVCRVVRYLQVVLLYASTYVLVSLSIDRYHAI

CYTOLOOP2 **TM4** **EXOLOOP2**
VYPMKFLQGEKQARVLIVIAWSLSFLFSIPTLIIFGKRTLSNGEVQCWAL

TM5 **CYTOLOOP3**
WPDDSYWTPYMTIVAFLVYFIPLTIISIMYGIVIRTIWIKSKTYETVISN

TM6
CSDGKLCSYYNRGLISKAKIKAIKYSIIILAFICCWSPYFLFDILDNFN

EXOLOOP3 **TM7**
LLPDTQERFYASVIIQNLPALNSAINPLIYCVFSSSISFPCRVIRLRQLQ

EAALMLCPQRENWKGTWPGVPSWALPR*

B_{short} (33 bp deletion in exon 3)

MPANFTEGSFDSSGTGQTLDSVPACTETVTFTEVVEGKEWGSF
 YYSFKTEQLITLWVLVFVFTIVGNSVVLFSTWRRKKKSRTFFVT
 QLAITDINWRFTGDFTAPDLVCRVVRYLQVVLLYASTYVLVSLSI
 DRYHAIVYPMKFLQGEKQARVLIVIAWSLSFLFSIPTLIIFGKRTLS
 NGEVQCWALWPDDSYWTPYMTIVAFLVYFIPLTIISIMYGIVIRTI
 WIKSKTYETVISNCSDGKLCSYYNRGLISKAKIKAIKYSIIILAFIC
CWSPYFLFDILDNFNLLPDTQERFYASVIIQNLPALNSAINPLIYC
VFSSSISFPCRVIRLRQLQEAAALMLCPQRENWKGTWPGVP
 SWALPR*

Figure 6B

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C

MPANFTEGSFDSSGTGQTLDSSPVACTETVTFTEVVEGKEWGSFYYSFKT
EQLITLWVLFVFTIVGNSVVLFSTWRRKKKSRMTFFVTQLAITV*

D

MPANFTEGSFDSSGTGQTLDSSPVACTETVTFTEVVEGKEWGSFYYSFKT
EQLITLWVLFVFTIVGNSVVLFSTWRRKKKSRMTFFVTQLAITGCAALRL
YLRPGVPQHRQIPCHRLPHEVPSRRKASQGPHCDRLEPVFSVLHSHPDHI
WEEDTVQR*

Figure 6C and 6D

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E

MPANFTEGSFDSSGTGQTLDSSPVACTETVTFTEVVEGKEWGSFYYSFKTEQLITLWVLF
VFTIVGNSVVLFSTWRKKKSRTFFVTQLAITDSFTGLVNILTDINWRFTGDFTAPDLVC
RVVRYLQKSKPGSSL*

F

MPANFTEGSFDSSGTGQTLDSSPVACTETVTFTEVVEGKEWGSFYYSFKTEQLITLWVLF
VFTIVGNSVVLFSTWRKKKSRTFFVTQLAITEKQARVLIVIAWSLSFLFSIPTLIIFG
KRTLSNGEVQCWLWPDDSYWTPYMTIVAFLVYFIPLTIISIMYGIVIRTIWIKSKYET
VISNCSDGKLCSSYNRGLISKAKIKAIKYSIIILAFICCWSPYFLFDILDNFNLLPDTQ
ERFYASVIIQNLPALNSAINPLIYCVFSSSISFPCREQRSQDSRMTFRERTERHEMQILS
KPEFI*

Figure 6E and 6F

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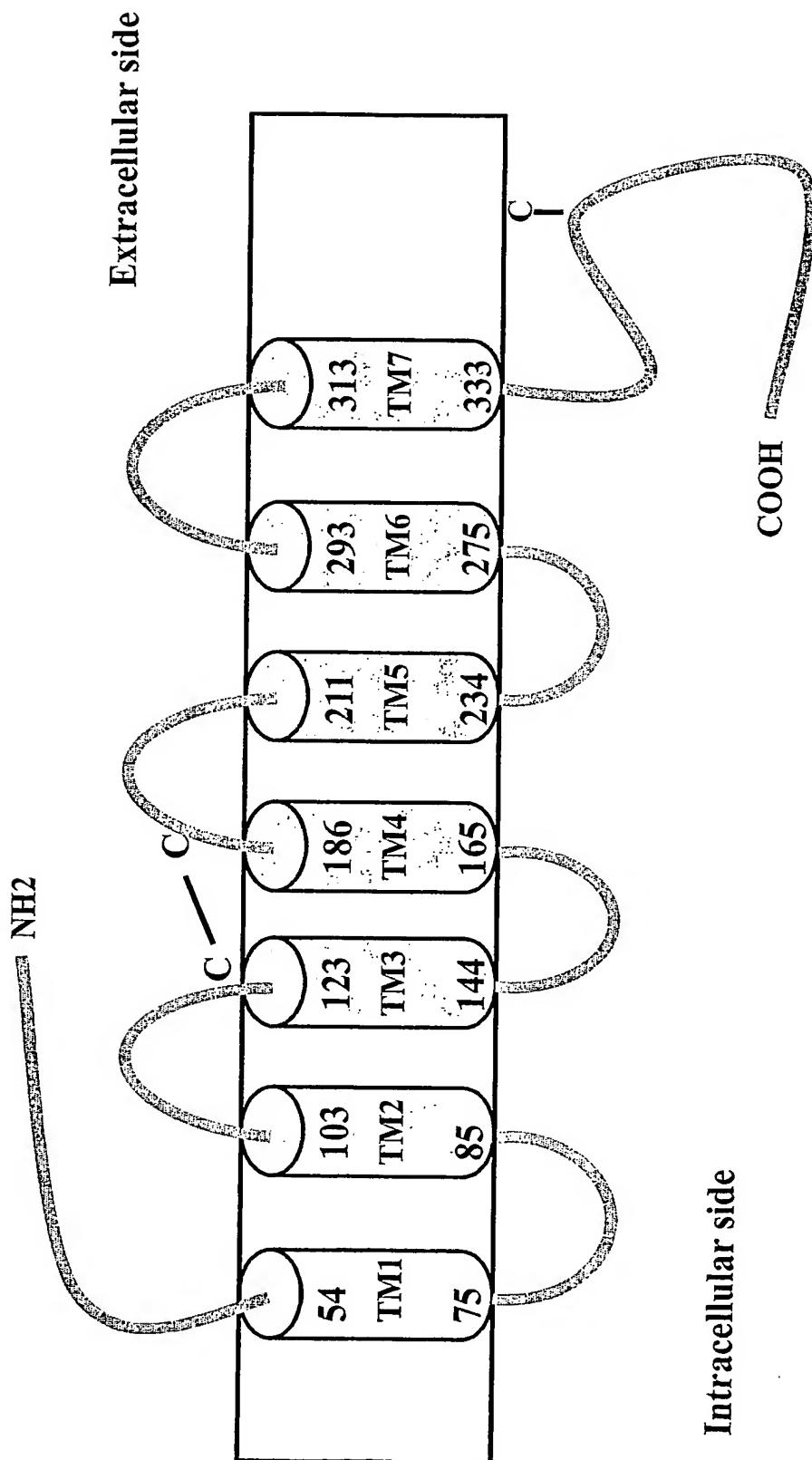


Fig. 7

101539565

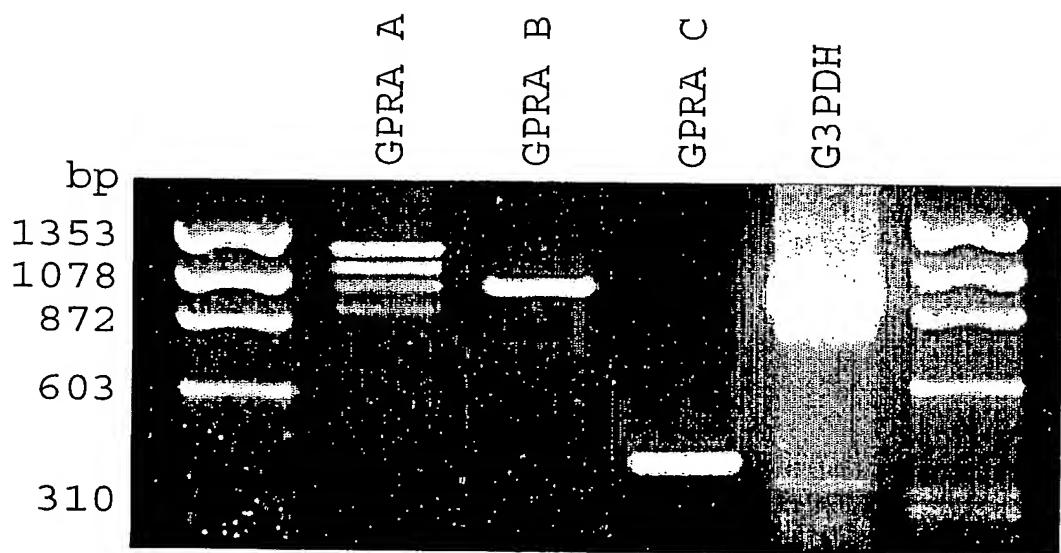


Figure 8

101539565

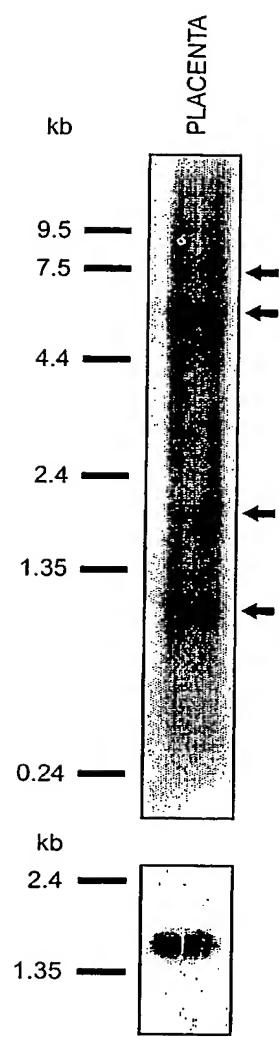


Figure 9

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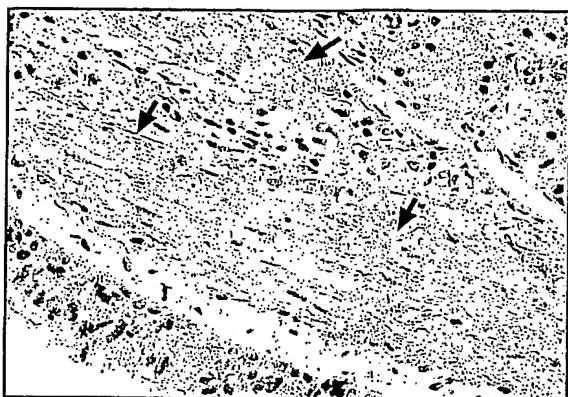
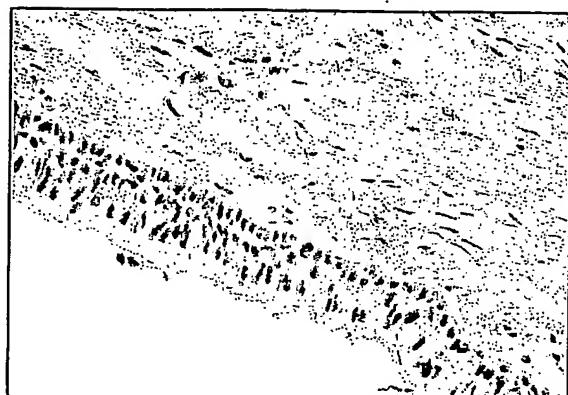
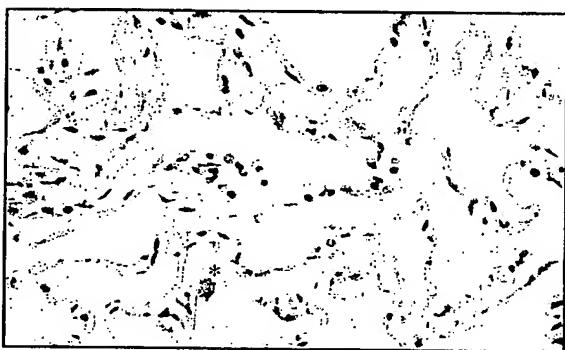
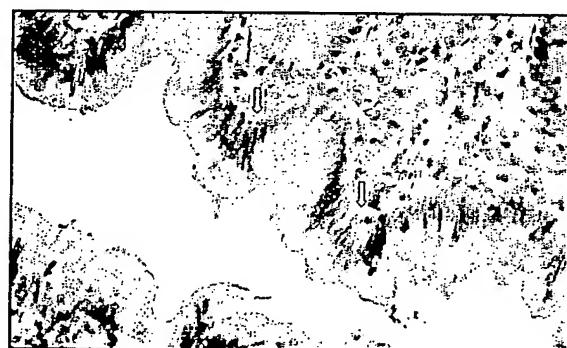
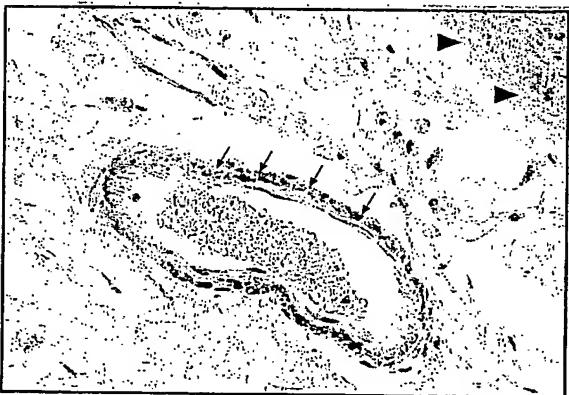
A.**B.****C.****D.****E.****F.**

Fig. 10A, Fig. 10B, Fig. 10C, Fig. 10D, Fig. 10E, Fig. 10F

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A.



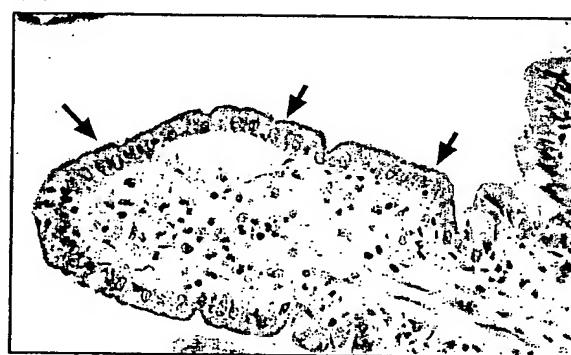
B.



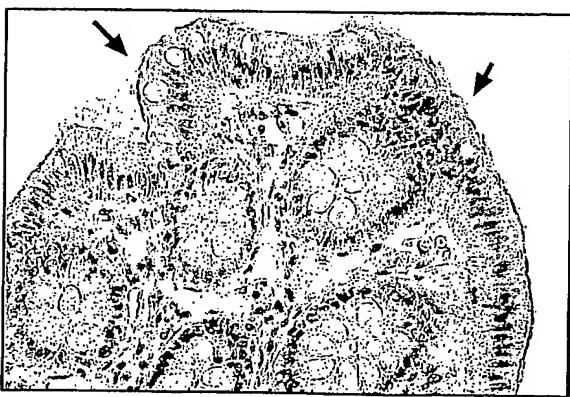
C.



D.



E.



F.

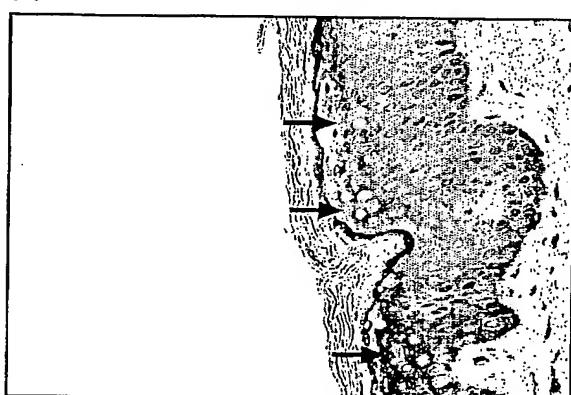


Fig. 11A, Fig. 11B, Fig. 11C, Fig. 11D, Fig. 11E, Fig. 11F

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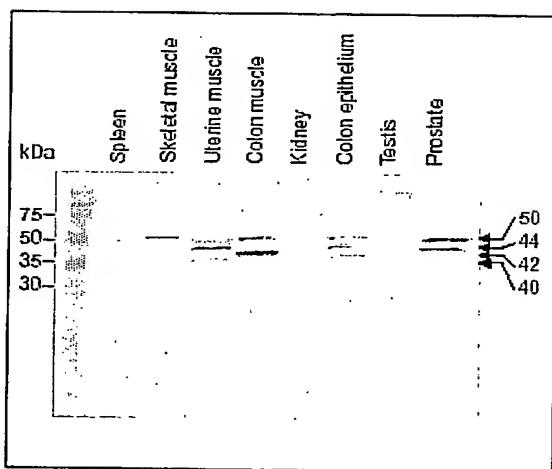
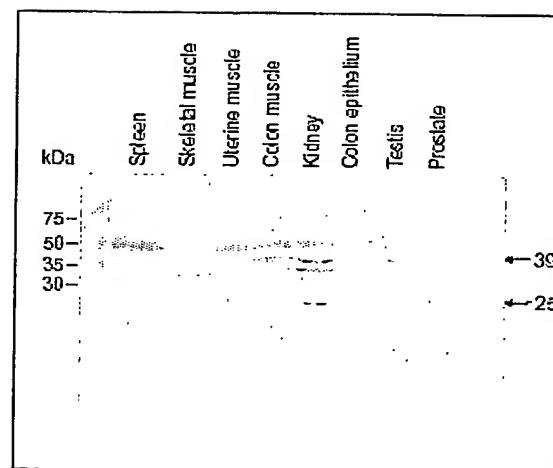
A.**B.**

Fig. 12A and 12B

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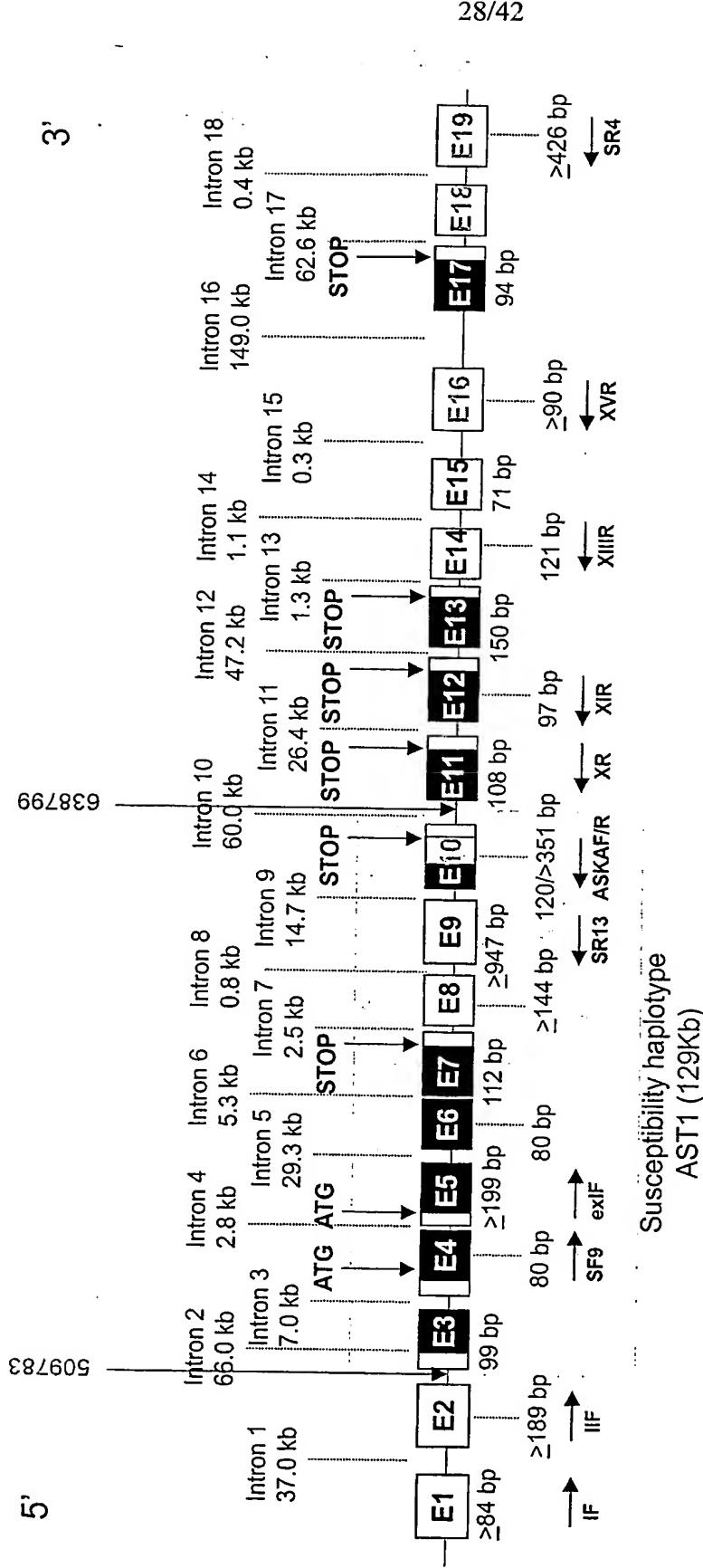


Fig. 13

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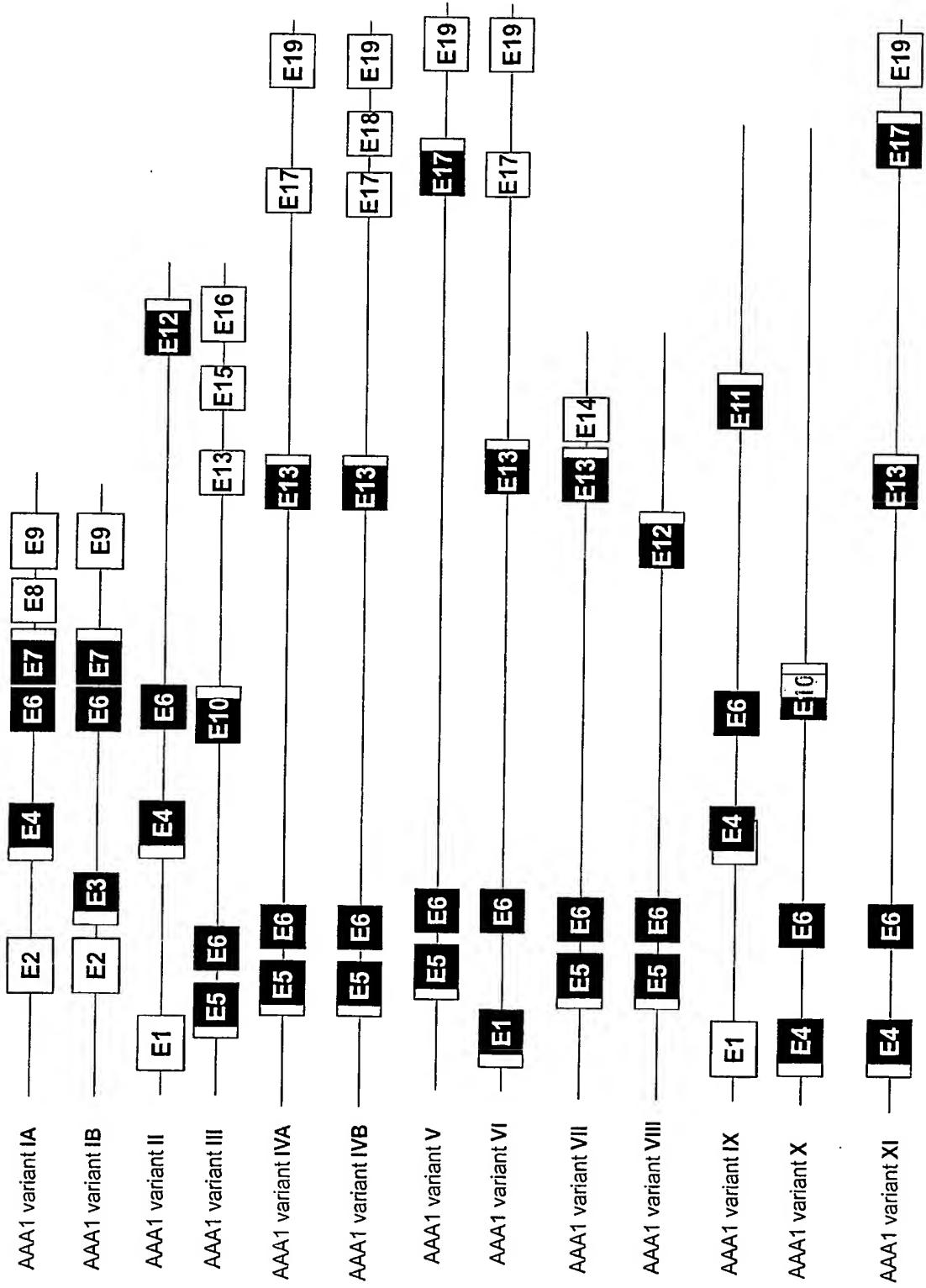


Fig. 14

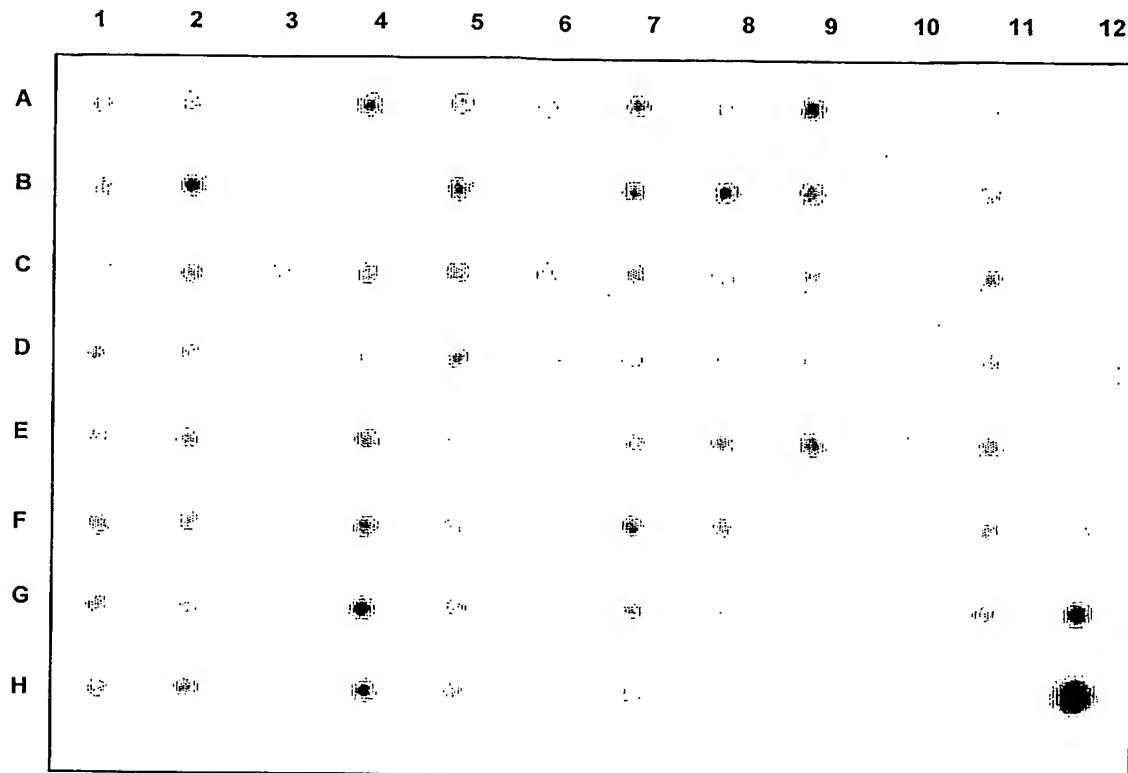
I	M Q C E A G A Y V R R N A G R Q F S H C N L H A H Q F L V R R K Q E L Q S V R W Y F B A G L W V K D T
II	M Q C E A G A Y V R R N A G R Q F S H C N L H A H Q F L V R R K Q V D I H C S K Q H V L K R F V F S P F N G L G T F L K N
III	M K K L K H R P A Y V R R N A G R Q F S H C N L H A H Q F L V R R K Q E N K D
*IV	M K K L K H R P A Y V R R N A G R Q F S H C N L H A H Q F L V R R K Q D W Q R L L P P L S C Q V G S P G C S A R K R S H V Q E H
V	M K K L K H R P A Y V R R N A G R Q F S H C N L H A H Q F L V R R K Q S K L L C I L Q I E C Y Q Q P Y K W R S R C F P S
VI	M P L D L M U E R L K T L G D I W K A Y V R R N A G R Q F S H C N L H A H Q F L V R R K Q D W Q R L L P P L S C Q V G S P G C S A R K R S H V Q E H
*VII	M K K L K H R P A Y V R R N A G R Q F S H C N L H A H Q F L V R R K Q D W Q R L L P P L S C Q V G S P G C S A R K R S H V Q E H
VIII	M K K L K H R P A Y V R R N A G R Q F S H C N L H A H Q F L V R R K Q V D I H C S K Q H V L K R F V F S P F N G L G T F L K N
IX	M Q C E A G A Y V R R N A G R Q F S H C N L H A H Q F L V R R K Q
X	M Q C E A G A Y V R R N A G R Q F S H C N L H A H Q F L V R R K Q E N K D
XI	M Q C E A G A Y V R R N A G R Q F S H C N L H A H Q F L V R R K Q D W Q R L L P P L S C Q V G S P G C S A R K R S H V Q E H

*) shared protein coding sequence, different 3'-untranslated regions

Figure 15

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	1	2	3	4	5	6	7	8	9	10	11	12
A	whole brain	cerebellum, left		heart	esophagus	colon, transverse	kidney	lung	liver	leukemia, HL-60	fetal brain	yeast total RNA
B	cerbral cortex	cerebellum, right	accumbens nucleus	nora	stomach	colon, descending	skeletal muscle	placenta	pancreas	HeLa S3	fetal heart	yeast rRNA
C	frontal lobe	corpus callosum	thalamus	atrium, left	duodenum	rectum	spleen	bladder	adrenal gland	leukemia, K-562	fetal kidney	E. coli rRNA
D	parietal lobe	amygdala		atrium, right	jejunum		thyroid	uterus	thyroid gland	leukemia, MOLT-4	fetal liver	E. coli DNA
E	occipital lobe	caudate nucleus	spinal cord	ventricle, left	ileum		peripheral blood leukocyte	prostate	salivary gland	Burkitt's lymphoma, Raji	total spleen	Poly (A)
F	temporal lobe	hippo-campus		ventricle, right	illicum		lymph node	testis		Burkitt's lymphoma, Daudi	fetal thymus	human C ₀ -1 DNA
G	p. g. of cerebrum cortex	medulla oblongata		latero-ventricular septum	appendix		bone marrow	ovary		colorectal adenocarcinoma, SW480	fetal lung	human DNA 100 ng
H	pans	putamen		apex of the heart	colon, ascending		trachea			lung carcinoma, A549		human DNA 500 ng

* paracentral gyrus

Figure 16

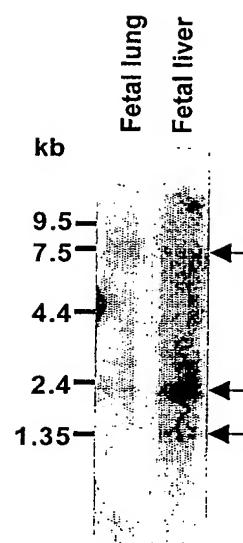


Figure 17

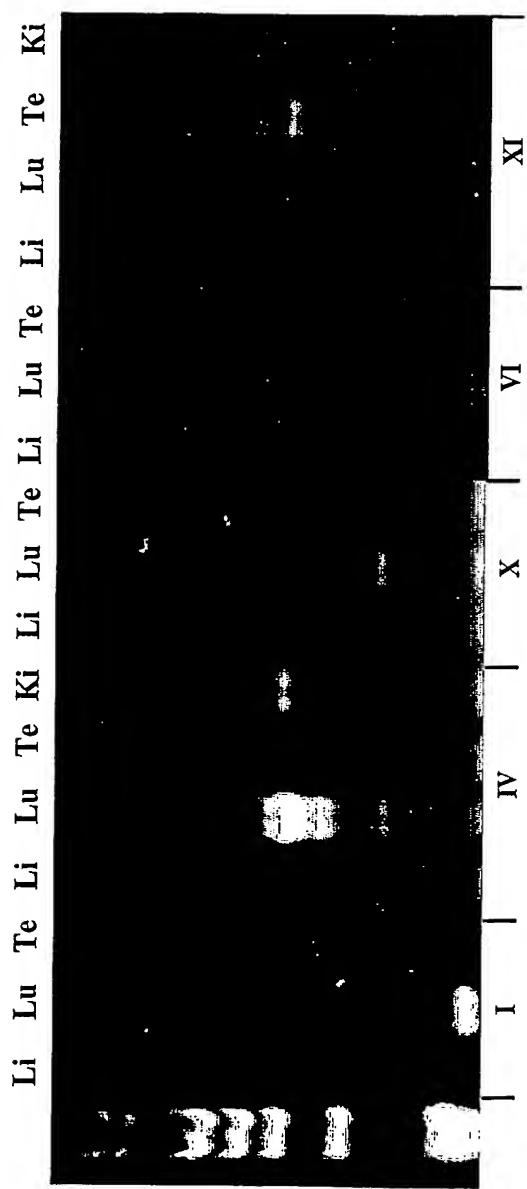


Figure 18

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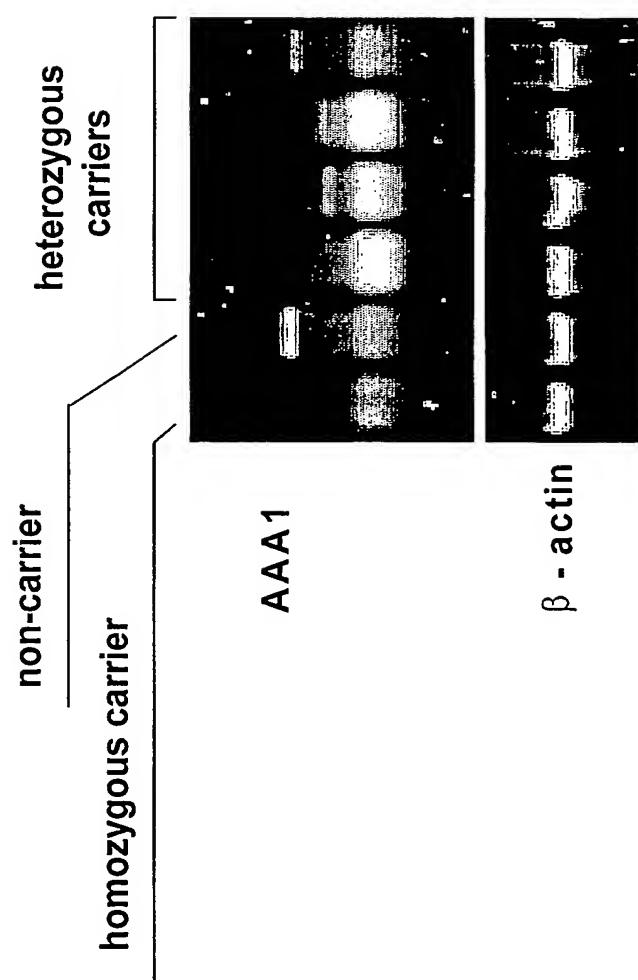


Figure 19

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Figure 20

I exons 2, 4, 6, 7, 8, 9 GB AY312364 SEQ ID NOS: 16 and 17

gacacagagaagaccactcccaccccccggagtgcgaaagggtgtgaaggggacagatcttttaac
catgcctgcccccttatacttgctgttcatagaattgcgaaactgaaagtgaccatgaggat
ccactggatggaggtaacttcttcttaagtggaggcataagatctggagtgacttctcc
ccagatttgtatacctgactctgtttcagcatccgctcccaaagaatgcagtgtgaa
M Q C E
gcaggagcttatgtgagaagaaaacgcaggagacagttcagtcactgcataatcttcatgcc
A G A Y V R R N A G R Q F S H C N L H A
catcagtttttgtgagaagaaaacaagagctccaatctgttagatgttattttgaagca
H Q F L V R R K Q E L Q S V R W Y F E A
ggctttgggtaaaggcacacctagaccctgaggactgaaaggcatggtgattattttggacaat
G L W V K D T
gggacatcaactctgtctatttgcataaaataagacttttgcacatgcacatggaggcag
gtcaaaagctccaggccaaactccaagttctgtatggggctctagccaaatggaaaggctct
tctccttcattgcctgactcttcaggactcttgcataactgcgaaagtggaaaatgagac
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ttgcataagagaatggaaacaatctgtctgtgattttagggacatactctggcagcaatat
ggaaatacagttcaatcctcattaacaaaacaggtatgaaatacatatttttagtaa
ggtgcctcactgtatgaaaaatccatttttccataatgttctgaaatgtcttag
cagtgcataagagacagcatgtcatcttcttagggactgtgttattgcattttcct
agggaagatcttcttaggtcacctgcctcgctaaagctctgaccaatctagctgc
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aagtgcacggccacatatgcacacgcataactgtgagggtattgcagtccttgc
ggttgcattgtataactggccagggtgttctatttccacattcttattaaatccct
acaggcagtttaggtatttagtgcacacacccctggcatagtcaccacatgcatt
agctccagataaactccagaaaaaagtccatccccacttctctcagtcgcctgccaac
gctggacacccctcaccacgcacagcagacagagaaaaagccctgggtttaagatcaaac
aaacacagctcaatttaggactctgtcacttcctgttactgggcacttgcgtgat
gtggttctcatctgtaaaacagagaaaaagatgattatctccaaatcttctatgttat
qttqaattaaataaqqtactctccatgaa

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Figure 20 (continued)

III exons 5, 6, 10a, 13, 15, 16

GB AY 312367 SEQ ID NOS: 22 and 23

tcttaggactcagaaaatatacgatgttagtaagagacaaacagacataacagataacacatac
aaagtgcctaccacatgctaaccactgctgcaggcactttctatagaagaactaattta
tcatcaccataaccctatgggtagatgatattttacaacacctccattttacagatgaag

M K
aaactgaagcatagacacctgcttatgtgagaagaaaacgcaggagacagttcagtcactgc
K L' K H R P A Y V R R N A G R Q F S H C
aatcttcatgccatcagttcttgagaagaaaacaagaaaacaaggactgaaatcca
N L H A H Q F L V R R K Q E N K D
cacaggaagggtggcagtgaactccacagacggacacctggacgcctcaacactcctggcctt
acctcccttgcgtgaacgtctcaagtttctgcgttcaggactggcaacgcctgcttc
cctctgagctgtcaagtaggaagtccgggctgcgtctgctagaaaagagaagtcatgtgcag
gagcactgaggcatcccagggtgtacactcttccacctagagcattccgtctctcatct
ctgccatgtgacgctggctttaacaaattaatcccaagtgcacagatatttatttct
tctgtacctaattgacactgagcaatcctctgtctgtaacctggtagtgtcatcttagaa
gtgaagacacaattaacacatggtcatttcttcattatatcgttqttact

IV exons 5, 6, 13, 17, 19

GB AY312368 SEQ ID NOS: 24 and 25

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Figure 20 (continued)

IVB exons: 5, 6, 13, 17, 18, 19

·SEQ ID NOS: 26 and 27

gtcttaggactcagaaaatatacatgttagtaagagcaaacagacataacagataacacataaaagtgcctaccacatgctaaccactgctgcaggcactttctatagaagaactaatttatacatcaccataaccctatgggttagatatatttacaacctccatttacagatgaag
M K
aaactgaagcatagacacctgttatgtgagaagaaaacgcaggagacagttcagtcactgc
K L K H R P A Y V R R N A G R Q F S H C
aatcttcatgccatcagttcttgtgagaagaaaacaagactggcaacgcctgttcct
N L H A H Q F L V R R K Q D W Q R L L P
cctctgagctgtcaagtaggaagtcgggctgctctgtctagaaaagagaagtcag
P L S C Q V G S P G C S A R K R S H V Q
gagcaactgaggcatcccaggtgtacactcttccacctagagcattccgtctctcatcct
E H
ctgccatgttagcaaaactgctatgcacccatcctcagctgcaaggattgaatgctatcaacaa
ccatacaagtggagaagcagatgcttccctagctgagcctcagctctgtttccctcagtc
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acaacttgggtgcacgcctgcctaaaaagaaaatactcaggaattgtctcataaagtgcctc
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tgagtccatcttccaaaggatctcgatttccctttaatgaaaataacattaaacacc
aaatataagcctcgctgtcccacatgcgtattgggacaagatgaaaacctgctccaggc
tactttggcagcagaactgaaaaaggctttccagatatactgatttctcatcgacag
ggttgcacagccctttatgttgcgttaatgacacccttggatctgaacaatacaca
ccaggacaattgtgtqacaaacttctacaaactqatatttctaatta

V exons 5, 6, 17, 19

GB AY 312369 SEQ ID NOS: 28 and 29

tcttaggactcagaaaatatacatgttagtaagagacaaacagacataacagataaacacata
aaagtgcctaccacatctaaccactgctgcaggcactttctatagaagaactaatttac
tcatcaccataaccctatgggtagatgatattttacaaccccttacatgtacatgtaa

M K
aaactgaagcatagacctgttatgtgagaagaaaacgcaggagacagttcagtcactgc
K L K H R P A Y V R R N A G R Q F S H C
aatcttcatgccatcagttcttgagaagaaaacaaagcaactgctatgcattcc
N L H A H Q F L V R R K Q S K L L C I L
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Q L Q G I E C Y Q Q P Y K W R S R C F P
agctgagcctcaggctttgatgaaattgctacaacttggatgcattccctaaaa
S
gaaataactcaggaattgtctcataaagtccctcacctactggcaaaaaacaagatgttctac
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tttcctttatgaaaataacattaaacaccaaataagcctcgatgtccacatcg
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tttttccagatatatgattctcatcgacagggttgcacagccctttattgttcgtg
taaatgacacccttggatctgaacaatacacaccaggacaattgtgtcaacagttctac
aaactgatatttctaatta

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Figure 20 (continued)

IB: exons 2, 3, 6, 7, 9

GB AY312365 SEQ ID NOS: 18 and 19

gacacagagaagaccactcccaccccccggagtgcaggtgtgaaggcacatcttt
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atccactggatggagttacttctttaagtgaggaggctaagatctgagttttcaca
tctctctgttagataaaaattccggctgggttccatctgtcagaagaactttctt
taatgtttcttaagtacaggtctgctgttatgtgagaagaaacgcaggagacagttc
C F L K Y R S A A Y V R R N A G R Q F
agtcaactgcaatcttcatgccatcagttcttgtgagaagaaaacaagagctccaatct
S H C N L H A H Q F L V R R K Q E L Q S
gttagatggtatttgaagcaggtcttggtaaaggacacccatagaccaggtaaggtca
V R W Y F E A G L W V K D T .
tggtagatttattggacaatgggacatcaactctgttattaaatgtgagaaaatgagacagg
ttgcactgagggtgttagccagacagactctcgacttggaaagtccatcttagatgttttgg
cataagagaatggaaacaatctgtctgtgatttagggacataactctgcagcaatatggg
aatacagttcaatccctattaacaaaacaggtatgaaatacatatttttagtaaggt
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ggcagttattaggtattgagtgctcacacacccctggcatagtcaccacatgcattagc
tccagataaactccagaaaaaagtccatccccacttctctcagctgcctgcacacgc
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gtttctcatctgtaaaacagagaaaagatgattatctccaaatcttctatgttatgtt
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II exons 1, 4, 6, 12

GB AY312366 SEQ ID NOS: 20 and 21

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gtttcagcatccgcttcccaaagaatgcagtgtgaagcaggagcttatgtgagaagaaca
M Q C E A G A Y V R R N
gcagggagacagttcagtcactgcaatcttcatgccatcagttcttgcataaaaaaa
A G R Q F S H C N L H A H Q F L V R R K
caagtggatatacactgttccaagcagcatgtgtgaaaagattgtctttccccattt
Q V D I H C S K Q H V L K R F V F S P F
aatggtcttggtaccttctcaaaaattgaccatatatga
N G L G T F L K N

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Figure 20 (continued)

VI exons 1, 6, 13, 17, 19

GB AY 312370 SEO ID NOS: 30 and 31

gctgtatgggtggaaaggagaatgagtcgttatgtgccttggacttgatgtggaaagacttaa
M P L D L M L E R L K
actttggggactactggaaaagcttatgtgagaagaaaacgcaggagacagttcagtccac
T L G D Y W K A Y V R R N A G R Q F S H
tgcaatcttcatgccatcgatcttgcgtgagaagaaaacaagactggcaacgcctgcctt
C N L H A H Q F L V R R K Q D W Q R L L
cctccctctgagctgtcaagtaggaagtcgggctgctctgcttagaaaagagaagtcatgtg
P P L S C Q V G S P G C S A R K R S H V
caggagcactgaggcatcccaggtgtgacactcttccacctagagcattccgtctctcat
Q E H
cctctgccatgttagcaaactgtatgcatttcagctgcaaggattgaatgtatcaa
caaccatacaagtggagaagcagatgttccttagctgagcctcaggctttgtatggaa
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tcctcacctactggaaaaacaagatgttctactcccaggttgactttcaagccccaa
gatgttgagtccagccatttcagatctcgatttcctttaatggaaaataacattaa
acaccaaataaagcctcgatgtcccatcgatgttgggacaagatgaaacctgcttc
caggctactttggcagcagaactggaaaaaggcttttccagatatatgatttctcatc
gacagggttgcacagccctttatgttcgtgtaaatgacacccttggatctgaacaat
acacaccaggacaattgtgtgcaacagttcataaaactqatatttctaatta

VII exons 5, 6, 13, 14

GB AY 312371 SEO ID NOS: 32 and 33

tcttaggactcagaaaatatacgatgttagtaagagacaacagacataacagataacacatac
aaagtgcctaccacatctaaccactgctgcagcactttctatagaagaactaattaa
tcatcaccataaccctatgggttagatgatattttacaacacctccatttacagatgaag

M K
aaactgaagcatagacctgc ttatgtgagaagaaaacgcaggggagacagttcagtcactgc
K L K H R P A Y V R R N A G R Q F S H C
aatcttcatgccatcagttcttgtagaagaaaacaagactggcaacgcctgcttcct
N L H A H Q F L V R R K Q D W Q R L L P
cctctgagctgtcaagtaggaagtccgggctgc tctgctagaaaagagaagtcatgtgcag
P L S C Q V G S P G C S A R K R S H V Q
gagcaactgaggcatcccagggtgtgacactcttccacctagagcattccgtctctcatcct
E H
ctgccatgtgccatgtttgaaccactagatttagagggctcaagcaatttcttggaaattt
actctgaattctacgttagaccatttcatgtgtataccctctgtgagtcaccctcaggta
gggacattt

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Figure 20 (continued)**VIII exons 5, 6, 12****GB AY 312372 SEQ ID NOS: 34 and 35**

tctaggactcagaaatatagatgttagtaagagcaaacagacataacagataacacatac
 aaagtgcctaccacatgctaaccactgctgcaggcacattctatagaagaactaattaa
 tcataccataaccctatgggttagatgatattttacaacctccatttacagatgaag
 M K
 aaactgaagcatagacacctgcttatgtgagaagaaaacgcaggagacagttcagtcactgc
 K L K H R P A Y V R R N A G R Q F S H C
 aatcttcatgccatcagttcttgcggatatacactgttccaag
 N L H A H Q F L V R R K Q V D I H C S K
 cagcatgtgtgaaaagatttgcattttcccccatttaatggcttggtaccccttcaaa
 Q H V L K R F V F S P F N G L G T F L K
 aattgaccatataatga
 N

IX exons 1, 4, 6, 11**GB AY 312373 SEQ ID NOS: 36 and 37**

gctgatgggagaaggagaatgagtgactctgtatgccttggacttgatgtggaaagacttaa
 gactttggggactactggaaaggagtgacttctcccccagattttgtatacctgactct
 gtttcagcatccgcttcccaaagaatgcagtgtgaagcaggagctatgtgagaagaaac
 M Q C E A G A Y V R R N
 gcagggagacagttcagtcactgcaatctcatgccatcagttcttgcggatatacactgc
 A G R Q F S H C N L H A H Q F L V R R K
 caagtttaggaaaacttcctacacccctttgtggatgtctctggactaatgactcc
 Q V
 aggcgagaccaccgttcatgactacttgcggatgtggact
 gagctact

X exons 4, 6, 10b**GB AY 321515 SEQ ID NOS: 38 and 39**

gagtgacttctccccagattttgtatacctgactctgtttcagcatccgcttcccaaaga
 atgcagtgtgaagcaggagcttatgtgagaagaaaacgcaggagacagttcagtcactgc
 M Q C E A G A Y V R R N A G R Q F S H C
 aatcttcatgccatcagttcttgcggatatacactgc
 N L H A H Q F L V R R K Q E N K D
 cacaggaagggtggcagtgaactccacagacggacccgtggacgcctcaacactcctggcctt
 acctcccttgcgtcaacgtctcaagttctgcgttcaggtatgtataggagggttatg
 agggcagagaattcctaagctcattagtaattgcattcagaaaatgtgcatttgcatt
 agctaattttcccaatatgagaagattggccctaccagaaaaagggaaatgatttgcatt
 aatgtgcgcaaaaatatgtttcttcttgcatt
 tcttatttagtccgcattttattgcattt

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Figure 20 (continued)**XI exons 4, 6, 13, 17, 19****GB AY 321516 SEQ ID NOS: 40 and 41**

gagtgacttctcccaagatttgtataacctgactctgtttcagcatccgcattccaaaga
atgcagtgtgaaggcaggagcttatgtgagaagaaacgcaggagacagttcagtcactgc
M Q C E A G A Y V R R N A G R Q F S H C
aatcttcatgccatcagttcttgtgagaagaaaacaagactggcaacgcctgcttcct
N L H A H Q F L V R R K Q D W Q R L L P
cctctgagctgtcaagttaggaagtccgggctgctctgctagaaagagaagtcatgtcag
P L S C Q V G S P G C S A R K R S H V Q
gaggactgaggcatcccaggtgtacactcttccacccatgagcattccgtctcatcct
E H
ctgccatgttagcaaactgctatgcatttcagctgcaaggattgaatgctatcaacaa
ccatacaagtggagaagcagatgttccctagctgagcctcaggcttttgatggattg
ctacaacttggcatgcctgcctaaaagaaaataactcaggaattgtctataaagtcc
tcacctaactggaaaaacaagatgttctactcccaggttgacttttcaagccccaaagat
gtttagtgcagccatttccaaggatctcgatttcccttaatggaaaataacattaaaca
ccaaatataagcctcgctgtcccacatgcgtattggggacaagatgaaacactgcttccag
gctactttggcagcagaactgaaaaaggcttttccagatatatgattctcatcgac
agggttgcacagccctttatgttgcgttaatgacacccttggatctgaacaataca
caccaggacaattgtgtcaacagttctacaaactgatattctaatta

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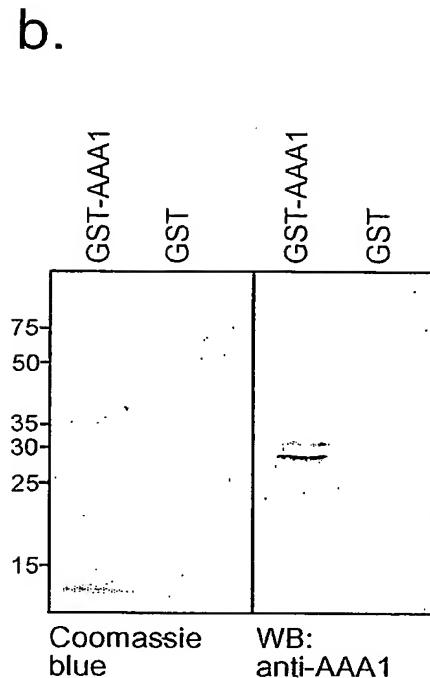
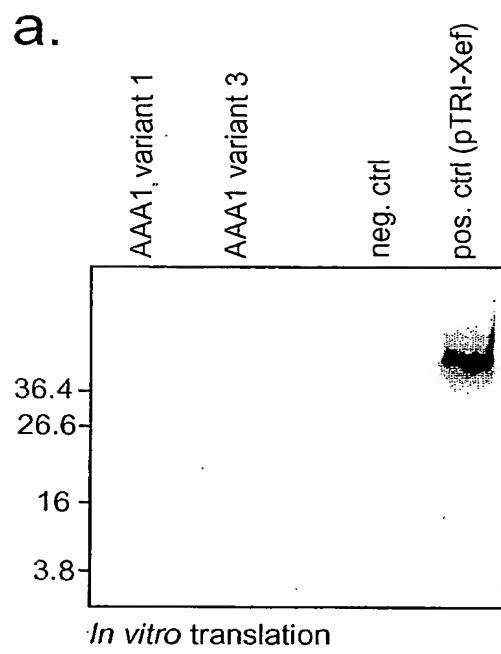


Fig. 21A and 21B